

Report

Decoding of Cytoplasmic Ca^{2+} Oscillations through the Spatial Signature Drives Gene Expression

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Summary

Cytoplasmic Ca^{2+} oscillations are a universal signaling mode that activates numerous cellular responses [1, 2]. Oscillations are considered the physiological mechanism of Ca^{2+} signaling because they occur at low levels of stimulus intensity [3]. Ca^{2+} oscillations are proposed to convey information in their amplitude and frequency, leading to activation of specific downstream targets [4–6]. Here, we report that the spatial Ca^{2+} gradient within the oscillation is key. Ca^{2+} oscillations in mast cells evoked over a range of agonist concentrations in the presence of external Ca^{2+} were indistinguishable from those in the absence of Ca^{2+} when plasmalemmal Ca^{2+} extrusion was suppressed. Nevertheless, only oscillations with accompanying Ca^{2+} entry through store-operated CRAC channels triggered gene expression. Increased cytoplasmic Ca^{2+} buffering prevented oscillations but not gene activation. Local Ca^{2+} influx and not global Ca^{2+} oscillations therefore drives gene expression at physiological levels of stimulation. Rather than serving to maintain Ca^{2+} oscillations by replenishing stores, we suggest that the role of oscillations might be to activate CRAC channels, thereby ensuring the generation of spatially restricted physiological Ca^{2+} signals driving gene activation. Furthermore, we show that the spatial profile of a Ca^{2+} oscillation provides a novel mechanism whereby a pleiotropic messenger specifically activates gene expression.

Results and Discussion

Since Prince and Berridge first proposed the existence of cytoplasmic Ca^{2+} oscillations in blowfly salivary gland [7], these oscillations induced by receptor activation have been found in virtually all cell types [8]. Ca^{2+} oscillations are thought to confer several advantages over a sustained bulk Ca^{2+} rise in activating Ca^{2+} -dependent responses [9]. These include increased sensitivity to the Ca^{2+} signal and circumvention both of the desensitization and excitotoxicity that can occur in response to a maintained Ca^{2+} signal. Furthermore, different amplitudes and/or frequencies of Ca^{2+} oscillation can recruit specific downstream Ca^{2+} -dependent targets [4–6, 10], providing a mechanism for selective responses to the promiscuous Ca^{2+} signal.

The diffusion of Ca^{2+} through an open Ca^{2+} channel in either the plasma membrane or an intracellular organelle results in the rapid build-up of a local microdomain of elevated Ca^{2+} [11]. Depending on the type of Ca^{2+} channel, the Ca^{2+}

concentration within a microdomain can reach tens of micromolar, several-fold higher than the bulk cytoplasmic Ca^{2+} rise [11, 12]. Ca^{2+} microdomains associated with voltage-gated Ca^{2+} channels activate neurotransmitter release [11], open colocalized Ca^{2+} -dependent K^{+} channels [13], and trigger gene transcription [14, 15], whereas microdomains that accompany the opening of store-operated Ca^{2+} release-activated Ca^{2+} (CRAC) channels activate the Ca^{2+} ATPase pump [16], adenylyl cyclase [17], endothelial nitric oxide synthase, and Ca^{2+} -dependent phospholipase A_2 [18]. Ca^{2+} microdomains arising from open InsP_3 receptors in the endoplasmic reticulum stimulate juxtaposed mitochondria to generate ATP [19]. Ca^{2+} oscillations arise from the opening of Ca^{2+} -permeable ion channels, so we hypothesized that the subcellular spatial profile of Ca^{2+} oscillations might play a role in activating selective downstream targets. If this is the case, then Ca^{2+} oscillations that are identical in amplitude and frequency should differentially activate downstream targets depending on the spatial location of the underlying Ca^{2+} channels. Here, we have generated Ca^{2+} oscillations that have identical amplitudes and frequencies but with distinct spatial profiles and we compared their abilities to activate gene expression.

To test the idea that the profile of a Ca^{2+} oscillation contained important information, we generated two kinds of InsP_3 -driven Ca^{2+} oscillation with distinct spatial signatures: one involved Ca^{2+} release from the endoplasmic reticulum without any Ca^{2+} influx, whereas the other comprised Ca^{2+} release followed by Ca^{2+} influx through store-operated CRAC channels. Only the latter scenario would result in significant elevation of the subplasmalemmal Ca^{2+} concentration.

Mast cells express cell-surface cysteinyl leukotriene type 1 receptors that couple to phospholipase C and which are activated by the proinflammatory molecule leukotriene C_4 (LTC_4) [20]. Stimulation with a submaximal concentration of LTC_4 (160 nM) in the presence of external Ca^{2+} evoked a series of repetitive Ca^{2+} oscillations that were maintained for at least 800 s (Figure 1A, representative of $n > 500$ cells with $\sim 98\%$ of cells responding). In the absence of external Ca^{2+} , oscillations ran down with time such that most cells failed to generate a Ca^{2+} signal after 400 s (Figure 1B, $n > 350$ cells, with virtually all cells responding). Hence Ca^{2+} influx is needed to sustain the Ca^{2+} oscillations. The Ca^{2+} influx pathway involves the CRAC channel because (1) in whole-cell patch clamp experiments, LTC_4 activates CRAC channels via store depletion [20] and (2) block of these channels with $1 \mu\text{M}$ Gd^{3+} or La^{3+} results in rundown of the oscillations in the presence of external Ca^{2+} and at a rate similar to that seen in the absence of external Ca^{2+} (Figure S1 available online; 56 cells). Ca^{2+} oscillations run down in Ca^{2+} -free solution because the Ca^{2+} that is released from the endoplasmic reticulum by InsP_3 is transported out of the cell, resulting in gradual depletion of the Ca^{2+} store. Of the plasma membrane Ca^{2+} removal mechanisms, the Ca^{2+} ATPase pump is particularly effective at clearing Ca^{2+} from the cytoplasm in mast cells [21]. After block of this pump with La^{3+} , stimulation with LTC_4 in the absence of external Ca^{2+} resulted in Ca^{2+} oscillations that were now sustained (Figure 1C; labeled 0 Ca^{2+} + La^{3+} , $n > 300$ cells, with

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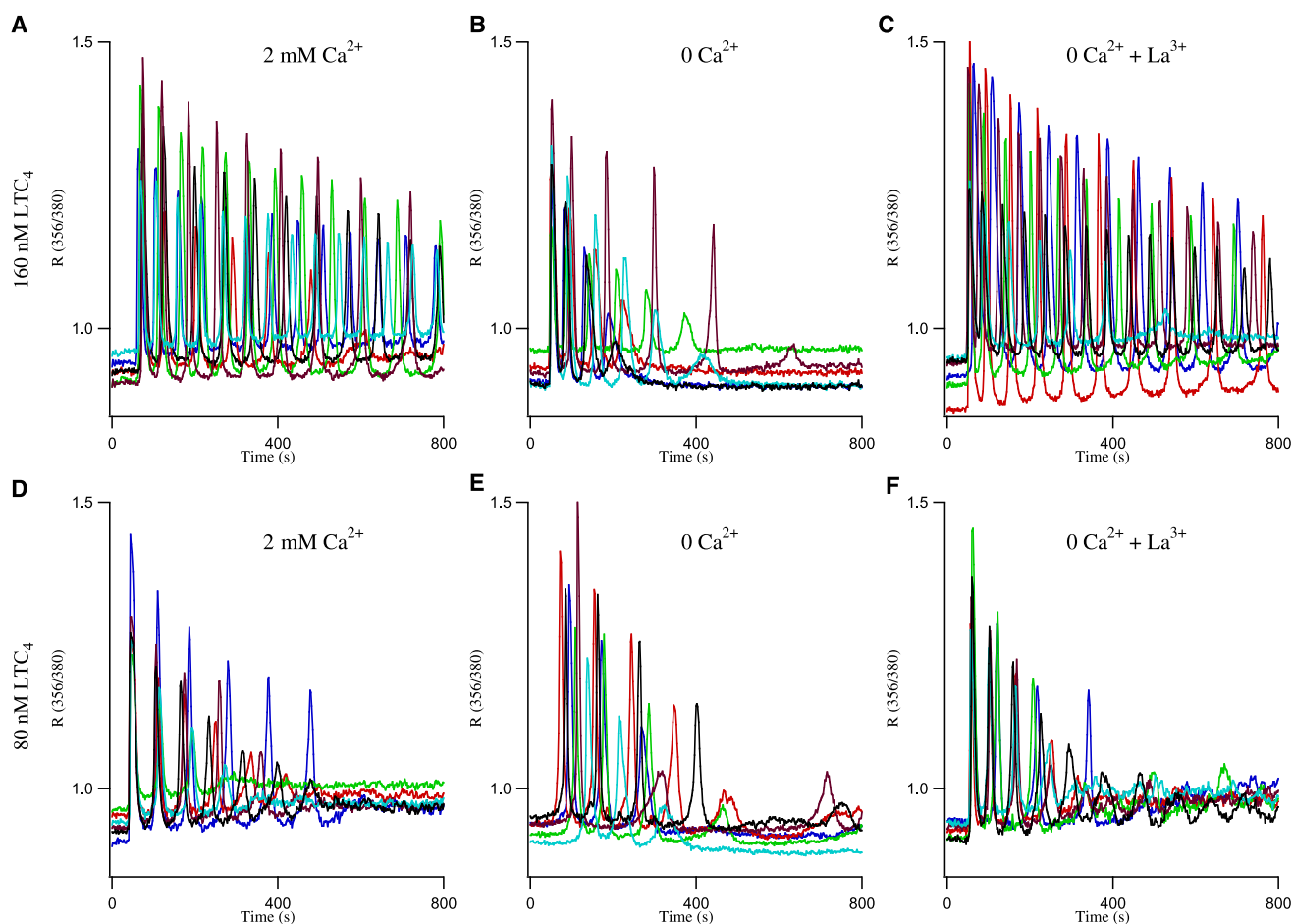


Figure 1. Cytoplasmic Oscillations Evoked by Low Levels of LTC₄

(A) Treatment with 160 nM LTC₄ evoked Ca²⁺ oscillations that were maintained in 2 mM external Ca²⁺. (B) Oscillations ran down in the absence of external Ca²⁺. (C) Oscillations were maintained in the absence of external Ca²⁺, if the plasma membrane Ca²⁺ATPase pump was blocked with La³⁺ (1 mM). (D–F) Ca²⁺ oscillations evoked by a lower concentration of LTC₄ (60 nM) showed a similar pattern. Each graph depicts different cells in a field of view. Different colors correspond to different cells. In each panel, LTC₄ was added at 30 s.

~all cells responding). Because repetitive Ca²⁺ oscillations can be supported in the absence of Ca²⁺ influx, Ca²⁺ entry does not directly contribute to their generation. Rather, in agreement with a previous study [22], it is needed to refill the stores and thus maintain the oscillatory mechanism that is driven by the intracellular milieu.

At lower agonist concentrations, the pattern of Ca²⁺ oscillations was indistinguishable between cells stimulated in 2 mM external Ca²⁺, 0 Ca²⁺, and 0 Ca²⁺ together with La³⁺ (Figures 1D–1F compares these responses in cells exposed to a low dose of LTC₄), although responses in 0 Ca²⁺ tended to run down slightly more quickly. We analyzed the amplitude and frequency of the Ca²⁺ oscillations for three different LTC₄ concentrations (Figure 2, each point represents >200 cells). The amplitude and frequency of the Ca²⁺ oscillations to LTC₄ were generally similar when evoked in either 2 mM Ca²⁺ or 0 Ca²⁺ and La³⁺, whereas responses in 0 Ca²⁺ were smaller and shorter lasting.

If distinct spatial Ca²⁺ gradients hidden within similar global Ca²⁺ oscillations encode important information, then one would predict these spatial profiles to differ in their ability to activate Ca²⁺-dependent responses. To test this, we compared the ability of the Ca²⁺ oscillations to activate expression of the gene *c-fos* because (1) it is an important component of the

transcription factor complex AP1 that regulates expression of chemokines in immune cells [23] and (2) Ca²⁺ influx through CRAC channels is a strong activator of *c-fos* transcription and translation [24]. Stimulation with LTC₄ for 12 min in the presence of external Ca²⁺ consistently activated *c-fos* expression over a range of agonist concentrations (Figure 3A). Strikingly, stimulation for the same time in the absence of Ca²⁺ failed to evoke any *c-fos* expression even when La³⁺ was present to maintain the Ca²⁺ oscillations (Figure 3A). This was the case over the range of LTC₄ concentrations tested (Figures 3B–3D). For identical amplitudes and frequencies, only Ca²⁺ oscillations in the presence of Ca²⁺ influx drove *c-fos* expression. Moreover, stimulation with 160 nM LTC₄ in Ca²⁺-free solution evoked 7 ± 1 Ca²⁺ oscillations in the 12 min period, whereas 80 nM LTC₄ in 2 mM Ca²⁺ generated only 3.5 ± 0.7 oscillations. Despite this, gene expression occurred only after challenge with LTC₄ in the presence of external Ca²⁺. The spatial Ca²⁺ gradient that arises from local Ca²⁺ influx through CRAC channels therefore has two important consequences: (1) it enables lower concentrations of an agonist to evoke a cellular response and (2) fewer Ca²⁺ oscillations are needed to activate gene expression, thus increasing the efficacy of excitation-transcription coupling.

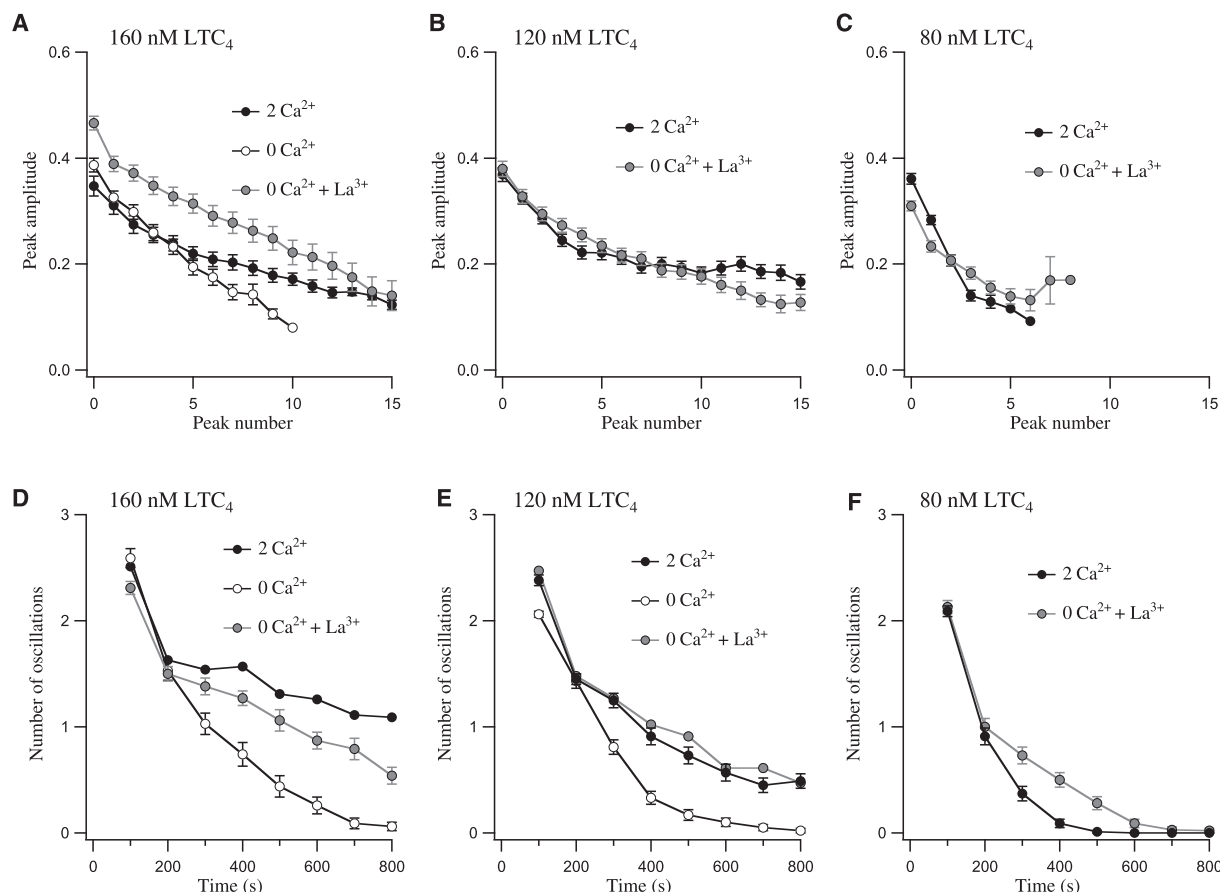


Figure 2. The Amplitude and Frequency of Ca^{2+} Oscillations to Three Different LTC_4 Concentrations Are Compared

(A–C) The amplitude of each Ca^{2+} oscillation against the oscillation number is plotted. The first Ca^{2+} oscillation is presented as peak 0, the second as peak 1, and so on.

(A) Response to 160 nM LTC_4 for the three different conditions shown.

(B and C) Response to 120 nM and 80 nM LTC_4 , respectively.

(D–F) The number of Ca^{2+} oscillations over an 800 s time frame is plotted. Oscillations were binned in 100 s intervals.

(D) Response to 160 nM LTC_4 .

(E and F) Response to 120 nM and 80 nM LTC_4 , respectively.

Data are presented as mean \pm SEM.

Stimulation of cysteinyl leukotriene type 1 receptors with LTC_4 activates CRAC channels [20] and a low concentration of Gd^{3+} (which is considered specific for CRAC channels) accelerated the decline of Ca^{2+} oscillations in 2 mM external Ca^{2+} (Figure S1). To confirm that Ca^{2+} entry through CRAC channels drives gene expression, we blocked the channels with the Synta compound and examined the effects on *c-fos* induction. As we have shown recently [25], the Synta compound is an effective inhibitor of CRAC channels. Stimulation with 160 nM LTC_4 in Ca^{2+} -free solution evoked a series of Ca^{2+} oscillations that ran down after ~ 7 min. Readmission of external Ca^{2+} resulted in a cytoplasmic Ca^{2+} rise resulting from Ca^{2+} influx through the open CRAC channels (Figure 4A). Pretreatment with 10 μM Synta compound substantially reduced this Ca^{2+} signal. Importantly, *c-fos* expression in response to LTC_4 (applied in the presence of 2 mM external Ca^{2+}) was prevented by the Synta compound (Figure 4B). Identical results were obtained with a lower concentration of LTC_4 (80 nM).

The finding that the amplitude and frequency of Ca^{2+} oscillations to LTC_4 were similar in 2 mM Ca^{2+} compared with 0 Ca^{2+} and La^{3+} suggests that Ca^{2+} entry and removal mechanisms

are in balance. CRAC channels cluster in punctate-like structures upon store depletion [26], but it is not known whether the plasma membrane Ca^{2+} pumps or sarco-endoplasmic reticulum Ca^{2+} ATPase pumps in peripheral endoplasmic reticulum colocalize here too. Nevertheless, it is likely that Ca^{2+} removal pathways are positioned such that they prevent a significant global Ca^{2+} increase after Ca^{2+} entry, while impacting little on the local subplasmalemmal Ca^{2+} rise in the vicinity of open CRAC channels. Clustering of removal mechanisms might be one effective way to ensure that local Ca^{2+} entry is impeded from eliciting a more global Ca^{2+} rise [27].

We considered the possibility that La^{3+} might enter cells and interfere with *c-fos* expression, thus complicating interpretation of the experiments in Figure 3 where gene expression is measured in 0 Ca^{2+} plus La^{3+} . Two arguments can be raised against this. First, we have previously shown that La^{3+} does not enter cells to any detectable level over the time course of the experiments described here [18]. Second, we triggered Ca^{2+} influx independent of store-operated Ca^{2+} entry to see whether La^{3+} inhibited the subsequent activation of gene expression. We used the Ca^{2+} ionophore ionomycin, which increases Ca^{2+} entry directly through its ionophore properties

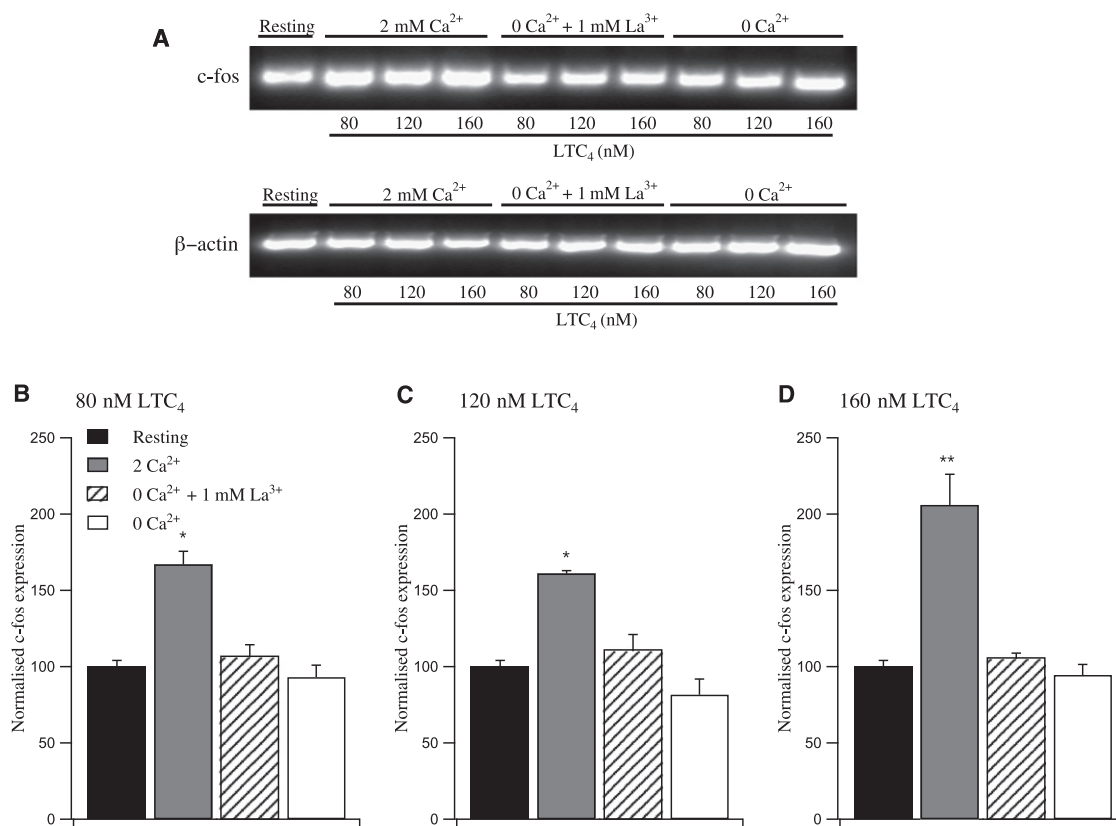


Figure 3. Gene Expression Occurs Only when Ca²⁺ Entry Accompanies the Oscillatory Signal

(A) *c-fos* expression is compared in cells exposed to different concentrations of LTC₄ in 2 mM Ca²⁺, 0 Ca²⁺ and La³⁺, or 0 Ca²⁺ alone.

(B–D) The extent of *c-fos* expression (normalized to control levels in nonstimulated cells) is compared for the different conditions after stimulation in 80 nM (B), 120 nM (C), and 160 nM LTC₄ (D). Over the concentration range tested, gene expression occurred only when accompanied by Ca²⁺ influx. Data are presented as mean ± SEM. *p < 0.01; **p < 0.001.

and indirectly through store depletion and subsequent activation of CRAC channels [28]. Because 1 mM La³⁺ completely abolishes store-operated Ca²⁺ entry (Figure 4C), we were able to isolate the Ca²⁺ influx because of direct ionophore transport by pretreating cells with this trivalent cation. The Ca²⁺ response to ionomycin was slightly reduced by 1 mM La³⁺ (Figure 4D) but *c-fos* expression was unaffected (Figure 4E). Hence La³⁺ does not interfere with Ca²⁺-induced gene expression.

Are global Ca²⁺ oscillations necessary for driving gene expression or is local Ca²⁺ influx sufficient? To distinguish between these possibilities, we increased cytoplasmic Ca²⁺ buffering by loading cells with the slow Ca²⁺ chelator EGTA. EGTA reduces the bulk Ca²⁺ rise but is too slow to interfere with the amplitude or extent of local Ca²⁺ entry [11, 12]. As we have shown previously, preincubating cells with EGTA-AM substantially reduced the Ca²⁺ rise that occurred after store-operated Ca²⁺ entry [18]. EGTA also abolished the global Ca²⁺ oscillations evoked by LTC₄ (63 cells) but had no inhibitory effect on *c-fos* expression in response to the same concentration of agonist (Figure 4F). We considered the possibility that the *c-fos* expression response might have already saturated after stimulation with thapsigargin in 2 mM Ca²⁺. If so, this could mask a potential difference in *c-fos* expression when compared with that induced after loading the cytoplasm with EGTA. However, a modest fall in the rate of Ca²⁺ entry produced a reduction in the extent of *c-fos* expression,

arguing against saturation of the response. Readmission of 0.25 mM and 0.5 mM external Ca²⁺ to cells pretreated with thapsigargin in Ca²⁺-free solution elicited a slower rate of Ca²⁺ influx (22.8% ± 6.7% and 48.9% ± 11.1% of that seen in 2 mM Ca²⁺ [18]). *c-fos* expression was likewise reduced to 38% ± 3.8% and 64% ± 5.1% (p < 0.01 in both cases) that seen in 2 mM Ca²⁺.

The Ca²⁺ chelator BAPTA is fast enough to intercept incoming Ca²⁺ and thus reduce the amplitude and spatial extent of local Ca²⁺ entry [11]. Loading the cytoplasm with BAPTA suppressed gene expression induced by LTC₄ (Figure 4F). Because BAPTA is able to restrict Ca²⁺ influx to within a few nanometers of the channel pore, these results demonstrate that local Ca²⁺ influx drives gene expression and that global Ca²⁺ oscillations are not essential to this process.

Our findings provide new insight into how cells decode the universal oscillatory Ca²⁺ signal. Rather than responding to the amplitude or frequency of the Ca²⁺ oscillations, we have discovered that the spatial profile of the oscillatory Ca²⁺ signal is the key trigger for gene expression. Abolition of global Ca²⁺ oscillations by increasing cytoplasmic Ca²⁺ buffering had no inhibitory effect on gene expression. Instead, local Ca²⁺ influx through CRAC channels was sufficient to drive *c-fos* transcription. The digital nature of the Ca²⁺ oscillations is therefore not the fundamental signaling unit used to drive excitation-transcription coupling at low levels of stimulus intensity, at least in the cell type we have used.

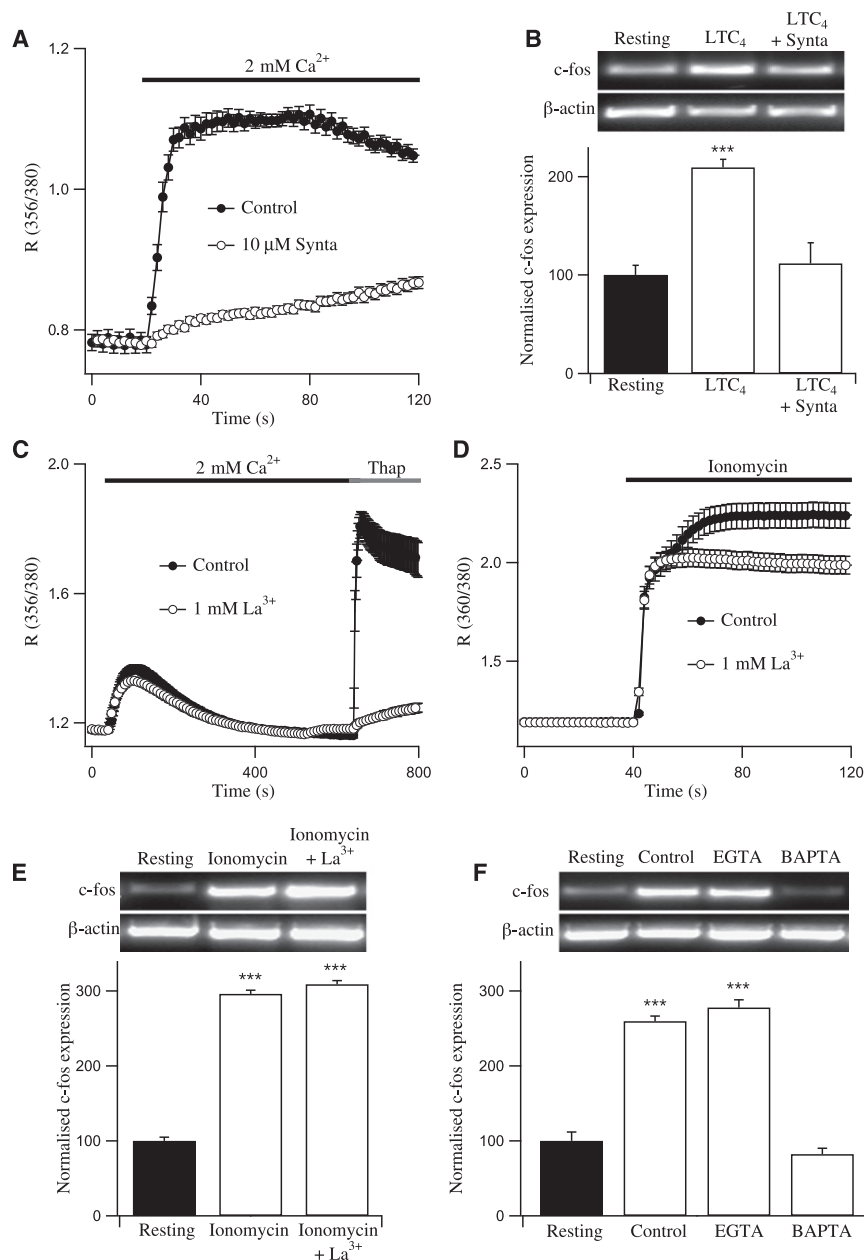


Figure 4. Gene Expression Can Occur in the Absence of Ca^{2+} Oscillations

(A) Cells were stimulated with LTC₄ (160 nM) in Ca^{2+} -free solution and once oscillations had stopped, 2 mM Ca^{2+} was applied. Store-operated Ca^{2+} entry occurred (filled circles, mean of >80 cells) and this was blocked by pretreatment with the Synta compound (10 μM , applied 5 min before Ca^{2+} was readmitted; >50 cells). Time 0 on the graph represents 60 s after cells had stopped oscillating.

(B) The Synta compound suppresses gene expression induced by LTC₄. Top: Gel shows *c-fos* induction by 160 nM LTC₄ and this is blocked by the Synta compound. Lower panel summarizes aggregate data from three such experiments.

(C) 1 mM La^{3+} abolishes store-operated Ca^{2+} influx. Thapsigargin (2 μM) was applied in 0 Ca^{2+} external solution to deplete the stores and then 2 mM external Ca^{2+} was readmitted as indicated.

(D) The Ca^{2+} signal evoked by 10 μM ionomycin was only partially reduced by La^{3+} .

(E) *c-fos* expression was induced by ionomycin and this was unaffected by La^{3+} .

(F) Loading cells with EGTA, which prevented the oscillatory response, had no effect on gene expression to LTC₄ whereas loading with BAPTA suppressed *c-fos* induction. Control refers to LTC₄ in 2 mM Ca^{2+} .

Data are presented as mean \pm SEM. *** $p < 0.0005$.

Although the importance of CRAC channels to physiological responses has been appreciated, it was thought that the role of the channels was simply to provide a “top-up” of Ca^{2+} stores, remedying the small loss of Ca^{2+} from the cell after each Ca^{2+} oscillation. Our data clearly reveal that the global Ca^{2+} rise associated with Ca^{2+} oscillations does not provide the Ca^{2+} that activates gene expression; rather, it is the spatially localized Ca^{2+} signal that arises from open CRAC channels that triggers the nuclear response. Hence CRAC channels do not simply support the signaling function of Ca^{2+} oscillations. On the contrary, it appears that the function of Ca^{2+} oscillations is to activate CRAC channels, which provides the important and physiological Ca^{2+} signal, at least for *c-fos* gene regulation.

The frequency of oscillatory Ca^{2+} signals can recruit different transcription factors to the nucleus [5], alter mitochondrial ATP production [6], and drive exocytosis [10]. Our results now show that important information is contained in the spatial signature

of the Ca^{2+} oscillation. In neurons, spatially restricted Ca^{2+} signals at the cell periphery can influence nuclear events: Ca^{2+} microdomains arising from L-type Ca^{2+} channels [15, 29] or NMDA receptors [30] activate gene expression independent of the bulk Ca^{2+} rise. Although these pioneering studies revealed the importance of local Ca^{2+} signals in driving excitation-transcription coupling, the stimuli used were 60–90 mM K^{+} pulses for tens of seconds or bursts of action potentials for 5 min. In our experiments, we have used a physiologically relevant trigger, namely a low dose of an agonist. Spatially restricted local Ca^{2+} entry can therefore trigger transcription after a stimulus of physiological intensity. Moreover, it is this spatial signature within a global Ca^{2+} oscillation, rather than the oscillation itself, that triggers *c-fos* gene expression. Such spatial decoding greatly increases the versatility of Ca^{2+} as a second messenger, enhances the efficacy of Ca^{2+} in promoting gene expression, and provides a novel means for generating specific Ca^{2+} -dependent cellular responses.

Experimental Procedures

Cell Culture

The mast cell line RBL-1 was bought from ATCC and cultured as previously described [25].

Single-Cell Ca^{2+} Imaging

RBL-1 cells were loaded with fura 5F as described [18]. Cells were excited at 356 and 380 nm (20 ms exposures) at 0.5 Hz and the ratio (R) calculated.

Standard external solution contained (in mM): NaCl 145, KCl 4.5, CaCl_2 2, MgCl_2 1, HEPES 10, D-glucose 10 (pH 7.2 with NaOH). In 0 Ca^{2+} solution, external Ca^{2+} was removed and 0.1 mM EGTA (pH 7.2 with NaOH) was added. 0 Ca^{2+} + La^{3+} was identical to 0 Ca^{2+} but 1 mM La^{3+} was added (and the pH returned to 7.2 with NaOH because EGTA can bind La^{3+}). LTC_4 was purchased from Axxora.

EGTA and BAPTA Loading

Cytoplasmic loading with EGTA or BAPTA was carried out by preincubating cells with EGTA-AM or BAPTA-AM (25 μM) for 40 min at $\sim 24^\circ\text{C}$ in the dark together with fura 5F (2 μM). Cells were then washed three times and allowed to recover for a further 15 min in the dark.

c-fos Expression

RT-PCR was carried out as described previously [24].

Supplemental Data

Supplemental Data include one figure and can be found with this article online at [http://www.cell.com/current-biology/supplemental/S0960-9822\(09\)00917-8](http://www.cell.com/current-biology/supplemental/S0960-9822(09)00917-8).

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